

Appl. No. : ~~unknown~~ 10|781,599
Filed : herewith

AMENDMENTS TO THE CLAIMS

1-41. (Canceled)

¹ ~~42.~~ (Currently amended) A method for detecting a non-B, non-C, non-G hepatitis virus gene wherein PCR is performed by using the oligonucleotides ~~as defined in claim 41~~ selected from the group consisting of a nucleotide sequence shown in SEQ ID NO: 57, a nucleotide sequence shown in SEQ ID NO: 60, and a nucleotide sequence shown in SEQ ID NO: 61 as primers.

² ~~43.~~ (Original) A method for detecting a non-B, non-C, non-G hepatitis virus gene wherein PCR is performed by using an oligonucleotide having a nucleotide sequence shown in SEQ ID NO: 57 and an oligonucleotide having a nucleotide sequence shown in SEQ ID NO: 60, or an oligonucleotide having a nucleotide sequence shown in SEQ ID NO: 57 and an oligonucleotide having a nucleotide sequence shown in SEQ ID NO: 61 as primers.

44-58. (Canceled)

³ ~~59.~~ (New) An isolated non-B, non-C, non-G hepatitis virus genome comprising:

- (a) a DNA having the nucleotide sequence shown in SEQ ID NO: 1 or 62, or
- (b) a DNA which hybridizes with the DNA of (a) under stringent conditions of 24 mM Tris-HCl, pH 7.5/2.4 mM EDTA/1M NaCl at 67°C.

⁴ ~~60.~~ (New) The genome according to claim ³ ~~59~~, which comprises the DNA having the nucleotide sequence shown in SEQ ID NO: 1 or 62.